

Last time: bacterial growth in chemostat

nutrient influx: $j_0 = n_0 \cdot \mu$
 μ : dilution rate
 n_0 : nutrient conc at inflow

$$\dot{S} = r(n)S - \mu S$$

$$\dot{n} = n_0\mu - n\mu - r(n)S/Y \leftarrow \text{biomass yield} = \frac{S}{n}$$

mass conservation: $\dot{S} + Y\dot{n} = \mu[(n_0 - n(t))Y - S(t)]$

for non-trivial steady state solⁿ ($S^* > 0, n^* > 0$)

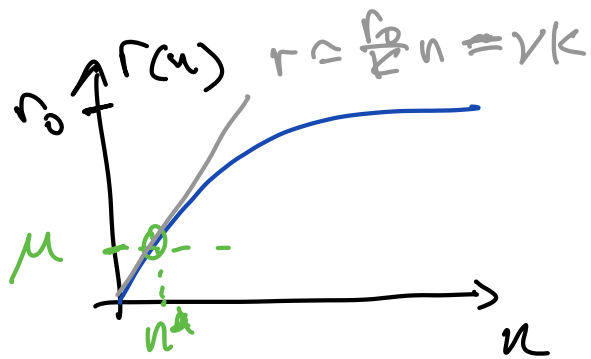
$$\dot{S} = 0 \text{ \& } \dot{n} = 0 \rightarrow S^* = (n_0 - n^*) \cdot Y \text{ (indep of } r(n))$$

value of n^* from $\mu = r(n^*)$

Monod growth law:

$$r(n) = r_0 \frac{n}{n+K}$$

$$\rightarrow \frac{n^*}{K} = \frac{\mu}{r_0 n^*} \xrightarrow{\mu \ll r_0} \frac{\mu}{r_0}$$



$$\frac{n^*}{n_0} = \frac{\mu K}{r_0 n_0} = \frac{\mu}{r_0 n_0} \equiv \eta \text{ (dimensionless parameter)}$$

$$S^* = \underbrace{n_0 Y}_{S_0} \left(1 - \frac{n^*}{n_0}\right) = S_0 \cdot (1 - \eta)$$

S_0 : max density attainable.

$\eta \rightarrow 1$: $S^* \rightarrow 0$ (washout)

$\eta \rightarrow 0$: $n^* \rightarrow 0, S^* \rightarrow S_0$ (complete conversion)

\Rightarrow Can estimate K by seeing how S^* depends on n_0 but maybe diff in practice due to proximity to washout

- Chemostat most stable when $\eta \ll 1$ (61)
- in this limit, $\frac{n^*}{n_0} \approx \eta \ll 1$.

$$p^* = (n_0 - n^*)\gamma \approx n_0\gamma = p_0$$

→ nutrient inflow mostly goes to biomass
(difficult to measure n^* , or $p_0 - p^*$)

- opposite limit $\eta \rightarrow 1$ (approaching wash out)

$$\frac{n^*}{n_0} \approx \eta \rightarrow 1, \quad (\text{e.g. reduce } n_0)$$

$$p^* = (n_0 - n^*)\gamma = (1 - \eta)p_0 \rightarrow 0$$

(Can estimate K , may be difficult to maintain)

b) Dynamics (relation to logistic growth model)

$$\dot{p} = (r(n) - \mu)p = (rn - \mu)p$$

$$\dot{n} = \mu(n_0 - n) - r(n)p/\gamma = \mu(n_0 - n) - vn p/\gamma.$$

[Compare to the damped predator-prey system (Sec A3)]

$$\begin{array}{l|l} \dot{p} = r p (1 - p/\tilde{p}) - b p q & \text{nutrient } (n) \leftrightarrow \text{prey } (p) \\ \dot{q} = c p q - \mu q & \text{cell } (p) \leftrightarrow \text{predator } (q) \\ & c \leftrightarrow v; b \leftrightarrow v/\gamma \end{array}$$

main difference:

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- prey replicates at rate $r \cdot (1 - p/\tilde{p})$
- nutrient injected at rate $\mu(n_0 - n)$

HW: In the limit $\eta \lesssim 1$ (near wash out)

dynamics converges to rapidly to

$$p(t) = [n_0 - n(t)] \cdot Y \quad (\text{mass conservation})$$

$$\begin{aligned} \text{Slow mode: } \frac{dp}{dt} &= (v n(t) - \mu) p = [v(n_0 - p(t)/Y) - \mu] p \\ &= [(v n_0 - \mu) - v p(t)/Y] p \end{aligned}$$

$$\frac{dp}{dt} = (r - p/\tilde{p}) \cdot p$$

→ recovers logistic eqn with effective "growth rate"

$$r = v n_0 - \mu = \mu \cdot (\tilde{r} - 1) \ll 1 \quad \text{for } \eta \lesssim 1$$

and "carrying capacity"

$$\tilde{p} = \frac{(v n_0 - \mu) Y}{v} = p_0 (\tilde{r} - 1) \ll p_0$$

(logistic eqn recovered here because it is leading order expansion in p when \tilde{p} is small)

II B. CR Model of Competition & Coexistence 63

1. Two-Species Interaction

a) 2-species growing on a single substrate
 chemostat with dilution rate μ

$$\dot{S}_1 = r_1(n) S_1 - \mu S_1$$

$$\dot{S}_2 = r_2(n) S_2 - \mu S_2$$

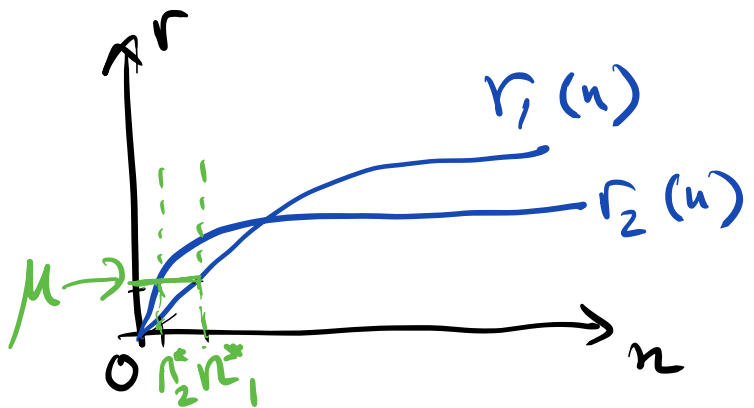
$$r_i(n) = r_i^0 \frac{n}{n + k_i}$$

$$\dot{n} = \mu(n_0 - n) - r_1(n) P_1 / Y - r_2(n) P_2 / Y$$

Steady state:

$$\mu = r_1(n^*), \quad \mu = r_2(n^*)$$

→ cannot be satisfied simultaneously unless $r_1(n) = r_2(n)$



Sol'n: $\mu = r_i(n^*)$, with $S_i \neq 0$, while $S_{j \neq i} = 0$.

→ i.e. only one species survives in steady state

HW: Surviving species is one with lower n_i

Γ approach: Assume one of the species goes extinct,

e.g. $S_2 = 0$.

check for stability for small $S_2 > 0$.

b) two species (of densities P_1, P_2) growing on two nutrients (of concentrations N_A, N_B)

- Chemostat with dilution rate μ .
- Nutrient influx ($j_A^0 = \mu N_A^0, j_B^0 = \mu N_B^0$)

Uptake of multiple nutrients:

- Substitutable. (e.g. glucose vs. glycerol)
- Essential (e.g. glucose + ammonium)

Substitutable nutrients:

- Many substrates co-utilized; GR approx additive

$$i.e. r_i(N_A, N_B) \approx r_i^0 \frac{N_A}{K_{iA}} + r_i^0 \frac{N_B}{K_{iB}}$$

$$\equiv \nu_{iA} N_A + \nu_{iB} N_B$$

[Hansen et al
MSB 2014]

- Some combo of substrates hierarchically utilized.

$$i.e., r_i(N_A, N_B) = \max \{ r_i(N_A), r_i(N_B) \}$$

essential nutrients: $r_i \approx \left(\frac{1}{\nu_{iA} N_A} + \frac{1}{\nu_{iB} N_B} \right)^{-1}$

→ will focus on substitutable, co-utilized nutrients in lectures; others in HW.

Dynamical equations:

$$\left. \begin{aligned} \dot{P}_1 &= (v_{1A}n_A + v_{1B}n_B)P_1 - \mu P_1 \\ \dot{P}_2 &= (v_{2A}n_A + v_{2B}n_B)P_2 - \mu P_2 \end{aligned} \right\}$$

$$\dot{n}_A = \mu(n_A^0 - n_A) - v_{1A}n_A P_1 / Y_A - v_{2A}n_A P_2 / Y_A$$

$$\dot{n}_B = \mu(n_B^0 - n_B) - v_{1B}n_B P_1 / Y_B - v_{2B}n_B P_2 / Y_B$$

assumption made: yield is species independent,
 Y_α can be scaled out $\left\{ \begin{array}{l} \text{but different substrate can contribute} \\ \text{quite differently to biomass,} \\ \text{e.g. glucose has 6C, glycerol only 3C} \end{array} \right.$
 $v_{i\alpha} / Y_\alpha = \tilde{v}_{i\alpha}$
 $n_\alpha Y_\alpha = \tilde{n}_\alpha$

* Steady state soln (for $P_1 \neq 0, P_2 \neq 0$)

$$\left. \begin{aligned} \dot{P}_1 = 0 & \quad v_{1A}n_A + v_{1B}n_B = \mu \\ \dot{P}_2 = 0 & \quad v_{2A}n_A + v_{2B}n_B = \mu \end{aligned} \right\} \begin{cases} \begin{bmatrix} v_{1A} & v_{1B} \\ v_{2A} & v_{2B} \end{bmatrix} \begin{bmatrix} n_A \\ n_B \end{bmatrix} = \begin{bmatrix} \mu \\ \mu \end{bmatrix} \end{cases}$$

Recall linear algebra

$$M \cdot \begin{bmatrix} x_1 \\ x_2 \end{bmatrix} = \begin{bmatrix} b_1 \\ b_2 \end{bmatrix} \rightarrow \begin{bmatrix} x_1 \\ x_2 \end{bmatrix} = M^{-1} \cdot \begin{bmatrix} b_1 \\ b_2 \end{bmatrix} \quad \left\{ \begin{array}{l} M = \begin{bmatrix} m_{11} & m_{12} \\ m_{21} & m_{22} \end{bmatrix} \\ M^{-1} = \frac{1}{\det M} \begin{bmatrix} m_{22} & -m_{12} \\ -m_{21} & m_{11} \end{bmatrix} \end{array} \right.$$

$$\begin{bmatrix} n_A^* \\ n_B^* \end{bmatrix} = \frac{1}{\det(V)} \begin{bmatrix} v_{2B} & -v_{1B} \\ -v_{1A} & v_{2A} \end{bmatrix} \begin{bmatrix} \mu \\ \mu \end{bmatrix} = \frac{\mu}{\det(V)} \begin{bmatrix} v_{2B} - v_{1B} \\ v_{1A} - v_{2A} \end{bmatrix} \sim 0 \left(\frac{\mu}{v} \right)$$

Note: nutrient levels set by v_{ij} and μ .
 not dependent on P_i , nor n_i^0 (cf. chemostat)

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P_1^*, P_2^* found from $\dot{n}_A=0, \dot{n}_B=0$

$$V_{1A} n_A^* P_1^* + V_{2A} n_A^* P_2^* = \mu (n_A^0 - n_A^*) Y_A$$

$$V_{1B} n_B^* P_1^* + V_{2B} n_B^* P_2^* = \mu (n_B^0 - n_B^*) Y_B$$

$$\underbrace{\begin{bmatrix} V_{1A} & V_{2A} \\ V_{1B} & V_{2B} \end{bmatrix}}_{V^T} \begin{bmatrix} P_1^* \\ P_2^* \end{bmatrix} = \mu \begin{bmatrix} \left(\frac{n_A^0 - n_A^*}{n_A^*}\right) Y_A \\ \left(\frac{n_B^0 - n_B^*}{n_B^*}\right) Y_B \end{bmatrix}$$

$$\begin{bmatrix} P_1^* \\ P_2^* \end{bmatrix} = \frac{\mu}{\det(V^T)} \begin{bmatrix} V_{2B} & -V_{2A} \\ -V_{1B} & V_{1A} \end{bmatrix} \begin{bmatrix} \left(\frac{n_A^0 - n_A^*}{n_A^*}\right) Y_A \\ \left(\frac{n_B^0 - n_B^*}{n_B^*}\right) Y_B \end{bmatrix} = \frac{\mu}{\det(V^T)} \begin{bmatrix} V_{2B} \frac{n_A^0 - n_A^*}{n_A^*} Y_A - V_{2A} \frac{n_B^0 - n_B^*}{n_B^*} Y_B \\ -V_{1B} \frac{n_A^0 - n_A^*}{n_A^*} Y_A + V_{1A} \frac{n_B^0 - n_B^*}{n_B^*} Y_B \end{bmatrix}$$

from above: $n_A^* = \mu (V_{2B} - V_{1B}) / \det(V)$ \rightarrow $1/n_A^* = \det(V) / [\mu (V_{2B} - V_{1B})]$
 $n_B^* = \mu (V_{1A} - V_{2A}) / \det(V)$ \rightarrow $1/n_B^* = \det(V) / [\mu (V_{1A} - V_{2A})]$

$$\begin{bmatrix} P_1^* \\ P_2^* \end{bmatrix} = \begin{bmatrix} \frac{(n_A^0 - n_A^*) Y_A}{1 - V_{1B}/V_{2B}} + \frac{(n_B^0 - n_B^*) Y_B}{1 - V_{1A}/V_{2A}} \\ \frac{(n_A^0 - n_A^*) Y_A}{1 - V_{2B}/V_{1B}} + \frac{(n_B^0 - n_B^*) Y_B}{1 - V_{2A}/V_{1A}} \end{bmatrix}; \det V = \det V^T$$

Note 1: $P_1^* + P_2^* = (n_A^0 - n_A^*) Y_A + (n_B^0 - n_B^*) Y_B$ — mass conservation

Note 2: if $\mu \rightarrow 0$, then $n_A^* \rightarrow 0, n_B^* \rightarrow 0$ (little nutrient left)

P_1^*, P_2^* depend on $(n_A^0 Y_A, n_B^0 Y_B)$ + V_{ij}

Note 3: if $n_A^0 - n_A^* \rightarrow 0, n_B^0 - n_B^* \rightarrow 0$ (i.e. $n_A^0 \rightarrow \frac{\mu (V_{2B} - V_{1B})}{V_{1A} V_{2B} - V_{1B} V_{2A}}$)
 then $P_1^* \rightarrow 0, P_2^* \rightarrow 0$ (washout limit of chemostat)

Goal: Understand dependence of Coexistence ($P_1 > 0, P_2 > 0$) (67)
 vs. dominance ($P_1 > 0, P_2 = 0$ or vice versa) or extinction ($P_1 = 0$)
 for diff environmental parameters ($j_A^0 = \mu n_A^0, j_B^0 = \mu n_B^0$)
 and genetic parameters (V_{ix}, M)

- find fixed points (if $P_i \leq 0$, then no coexistence)
- if $P_A > 0, P_B > 0$ exist, determine stability
 - unstable in one direction: phase transition (multi-modality)
 - Stable = Coexistence occurs
 - basin of attraction?

A simple limit (weak interaction case):

Specialist (n_A) → P₁ V_{1A} ≫ V_{2A} → Coexistence.
 vs
 Specialist (n_B) → P₂ V_{2B} ≫ V_{1B} (if not washed out)

Q: as interaction is turned on,
 to what extent is coexistence stable?

Specialist ① (n_A) → P₁ V_{2B} ≫ V<sub>1A}, V_{1B} ??
 vs
 Generalist ② (n_B) → P₂ V_{1B} << V_{1A}, V_{1B} ?}</sub>

Generalist (n_A) → P₁
 vs
 Generalist (n_B) → P₂ all V_{ix} comparable ??

→ General analysis of stability (around $P_1^*, P_2^*, n_A^*, n_B^*$)

4x4 matrix - not intuitive

→ Short-cut:

effective dynamics of n_A, n_B (Tilman)